

Gene Section

Review

PCSK6 (proprotein convertase subtilisin/kexin type 6)

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Abstract

Review on PCSK6, with data on DNA/RNA, on the protein encoded and where the gene is implicated.

Identity

Other names: PACE4, SPC4

HGNC (Hugo): PCSK6

Location: 15q26.3

DNA/RNA

Description

This gene can be found on chromosome 15 at location: 101844134-102029873.

Transcription

The DNA sequence contains 23 exons and the transcript length: 4236 bps translated to a 968 residues protein.

Protein

Description

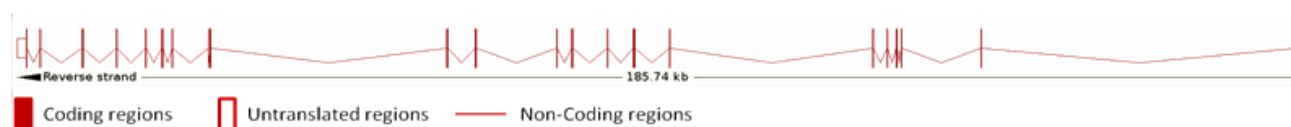
PCSK6 is a member of the family of subtilisin-like proprotein convertases (PCs) that process precursor

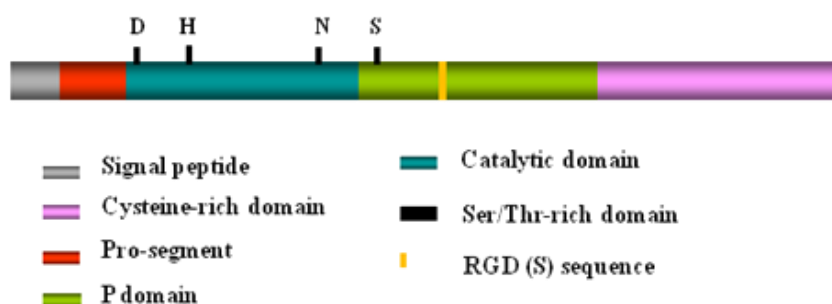
proteins at their paired basic amino acid processing sites (RXR).

This protease undergoes an initial autocatalytic processing event in the endoplasmic reticulum (ER) to generate a heterodimer which exits the ER and sorts to the trans-Golgi network (TGN). In the TGN a second autocatalytic cleavage takes place in order to release an active PCSK6.

Expression

PCSK6 is expressed in many tissues and encoded by different alternatively spliced mRNAs: the PACE4A-I form is secreted and expressed in heart, brain, placenta, lung, skeletal muscle, kidney, pancreas, but at comparatively higher levels in the liver. Isoform PACE4A-II is secreted and at least expressed in placenta. Isoform PACE4B is secreted and was only found in the embryonic kidney cell line from which it was isolated. Isoforms PACE4C and PACE4D are located in ER (not secreted, remain probably inactive unprocessed form in endoplasmic reticulum) and expressed in placenta too. Isoforms PACE4E-I and PACE4E-II are probably retained intracellularly through a hydrophobic cluster in their C-terminus, and expressed in cerebellum (E-I and -II), placenta and pituitary (only E-I). The PACE4CS was also found to probably lack enzymatic activity.





Localisation

PCSK6 is activated predominantly at the cell surface and able to process its substrates within the extracellular matrix and cell surface.

Function

PCSK6 induce protein precursors cleavage at the R-X-(K/R)-R motif to release mature proteins. PCSK6 substrates include enzymes (ADAMTS-4 and ADAMTS-5), HIV accessory protein Vpr, NODAL, TGF-beta related proteins, proalbumin and von Willebrand factor. This protein is thought to play a role in tumor progression and left-right patterning.

Homology

The PCSK6 catalytic domain has a high percentage of homology with those of the other PCs: 42% between PCSK6 and Furin.

Implicated in

Breast cancer

Higher expression of PCSK6 has been detected in human breast cancer. There was a correlation between the amount of PACE4 gene product and estrogen receptor content.

Prostate cancer

A recent study demonstrated that PCSK6 was highly overexpressed in prostate cancer tissues.

Skin cancer

Transgenic expression of PACE4 in keratinocytes of the epidermal basal layer was more susceptible to chemical carcinogenesis, evidenced by increased tumor multiplicity, tumor volume, and metastasis.

Association with degree of handedness

There is a significant association with a VNTR polymorphism in PCSK6 and handedness.

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